



CASE DB13NP

CERTIFICATE OF MAILING

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Keith R. Lange
Type or print name


Signature

December 4, 2002
Date

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

IN RE APPLICATION OF
STARLING ET AL.

APPLICATION NO: 09/745,605

FILED: DECEMBER 22, 2000

FOR: NOVEL IMMUNOGLOBULIN SUPERFAMILY MEMBERS OF APEX-
1, APEX-2 AND APEX-3 AND USES THEREOF

Art Unit: 1644

Examiner: Haddad, Maher M.

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Assistant Commissioner for Patents
Washington, D.C. 20231

**DECLARATION OF PRIOR INVENTION IN
THE UNITED STATES TO OVERCOME A REFERENCE UNDER 37 C.F.R. § 1.131**

Sir:

1. We, Gary C. Starling and Joshua N. Finger, both citizens of the United States, residing respectively at 11 Gallo Court, Lawrenceville, NJ 08648 and 197 North Union Street, Apt. B, Lambertville, NJ 08530 are joint inventors of the above-identified application.
2. At the time of the invention thereof we were working for Bristol-Myers Squibb Company, assignee of the present application. We submit this declaration to establish completion of the invention set forth in this application in the United States at a date prior to December 9, 1999, i.e. the publication date of WO 99/63088 to Baker et al. (hereinafter the '088 publication), which was cited by the Examiner in an Office Action mailed June 5, 2002.

3. From the documents submitted herewith and as set forth hereinbelow, it can be seen that the invention was completed in the United States before December 9, 1999, the publication date of the '088 publication. Completion of the invention prior to December 9, 1999 is shown by conception and actual reduction to practice of the invention as evidenced by the cloning and sequencing of the APEX-1 gene (hereinafter "APEX-1"), which is also referred to in Exhibit A as DCS4.
4. To establish conception and reduction to practice, i.e. completion of the invention at a date prior to December 9, 1999, the following documents are submitted as evidence:
 - a. Bristol-Myers Squibb Notebook No. 42973 assigned and completed prior to December 9, 1999 (Exhibit A), pages 42973-103 through 42973-106, 42973-112 through 42973-114, 42973-127 through 42973-129 and page 42973-158. These pages show the full-length cloning of APEX-1 and set forth the nucleotide and amino acid sequences of APEX-1, which correspond to SEQ ID NOS. 1 and 4, respectively, in the present application. The full length cDNA sequence and amino acid translation of APEX-1 is shown on page 42973-158. These notebook records evidence conception and actual reduction to practice of the complete invention prior to December 9, 1999.
5. The materials submitted herewith establish that the invention as claimed was completed, i.e. conceived and reduced to practice, at a date prior to December 9, 1999, the publication date of the '088 publication.

6. This declaration is submitted in a response to a non-Final Office Action dated June 5, 2002 and is therefore believed to be timely filed.

7. We hereby declare that all statements made herein of our own knowledge are true and that all statements made on information or belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and like so made are punishable by fine or imprisonment or both under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

DATED: _____

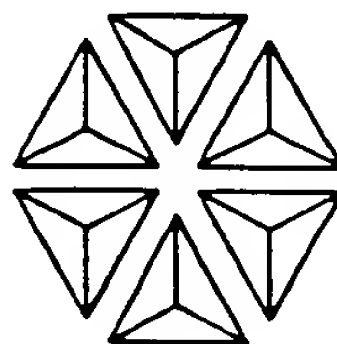
Gary C. Starling

DATED: _____

Joshua N. Finger

EXHIBIT A

BRISTOL-MYERS SQUIBB PHARMACEUTICAL RESEARCH INSTITUTE



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BRISTOL-MYERS SQUIBB

NOTEBOOK Nº 42973

Assigned to Joshua N. Fung

Department Name _____

Department Number _____

Date Assigned _____

Date Completed _____

Pages Completed from 001 to 200

Continued from Notebook Number _____

Continued in Notebook Number _____

This notebook cannot be transferred to another person

TABLE OF CONTENTS



PROJECT OR EXPERIMENT NO.	PRODUCT OR SUBSTANCE	STUDY PERFORMED OR OBJECTIVE	PAGES
		Construction of Dendritic substrate panel	090
		Hybridization of Southern	091
		Colonies from Stem selection - MAN's	092
		"	093
		cDNA synthesis	094
		Adoptive hybridization	095
		Dendritic clones	096
		"	097
		"	098
		Dendritic adoption hybridization	099
		"	100
		"	101
		BLANK	102
	DCS4 (Apex1)	DCS4 (clone #4) and Clone # B2 isolation	103
		"	104
		"	105
		BLANK Southern blot	106
		BLANK	107
		BLANK	108
		RNA isolation, THP1, NIH3T3, BTAB	109
		Northern hybrid of cell lines	110
		"	111
		DCS4 3'-RACE	112
		"	113
		"	114
		Northern blot of cell lines	115
		DCS4 3'-RACE	116
		"	117
		DCS4 northern analysis - MAN's	118

TABLE OF CONTENTS

PROJECT OR EXPERIMENT NO.	PRODUCT OR SUBSTANCE	STUDY PERFORMED OR OBJECTIVE	PAGES
		DCS4 MTN blots	119
		"	120
		DCS4 3'-RACE cloning	121
		" transformation	122
		"	123
		RT-PCR in cell lines - DCS4	124
		"	125
		"	126
		Full-length cloning DCS4	127
		RT-PCR using JNF6/7 - DCS4	128
WP761 Library		"	129
		RNA isolation from activated PBSTs (PHA)	130
		cDNA synthesis - WP761 library	131
		"	132
		"	133
		RsaI digestion - WP761 library	134
		"	135
		Hybridization	
		Adapter ligations - WP761 library	136
		"	137
		"	138
		DCS4 RT-PCR - cell lines	139
		"	140
		Ligation efficiency - WP761 library	141
		FLDCS4 minipreps	142
		Genomic Southern	143
		DCS4 JNF6/7 probe prep	144
		"	145
		Genomic PCR using primer pairs	146
		"	147

DATE

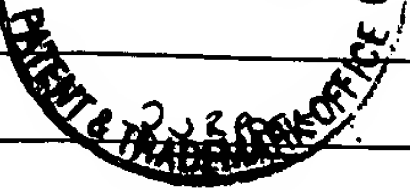
PROJ. NO.

EXPT. NO.

SUBJECT

Isolation of Clone #4 and Clone #82

DEC 08 2002



To isolate the cDNA inserts from clone #4 and clone #82 from plasmid DNA for use as probes.

5

Materials and Methods

a) Clone #4 0.5ul
 React III 0.1ul
 EcoRI 1ul
 dH₂O 3ul
 10ul

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b) Clone #82 19ul
 React II 3ul
 PstI 1.5ul
 HindIII 1.5ul
 dH₂O 5.0ul
 30.0ul

15

20

RESULTS:

Lane 1: 100 Marker
 Lane 2: Clone #4
 Lane 3: BLANK LANE
 Lane 4: BLANK LANE
 Lane 5: Clone #82
 Lane 6: Clone #82

25

30

Discussion: Clone #4 must have a mutated EcoRI site, cut again with HindIII and PstI. Go ahead and isolate clone #82 insert and analyze.

35

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SUBJECT



Reagent of Clone #4 using Pst I and Hind III

To drop out insert of clone #4 using enzymes other than EcoRI

5

Materials and Methods

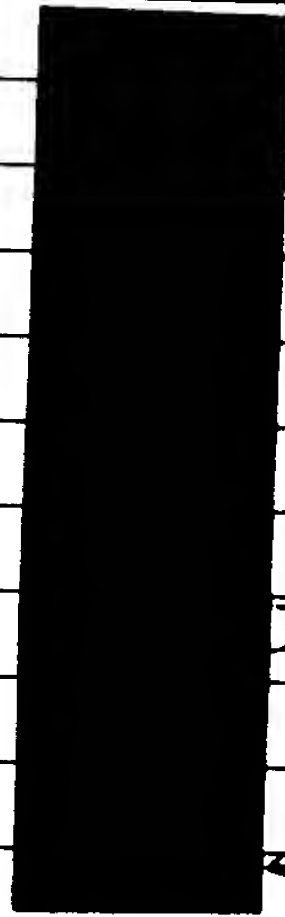
10

Clone #4	1.5 μ l
Pst II	3 μ l
Pst I	1.5 μ l
Hind III	1.5 μ l
dH ₂ O	9.0 μ l
	30.0 μ l

15 Results

Lane 1 ϕ marker
 Lane 2 H/P digest of DSS4
 Lane 3 H/P digest of DSS4

20



← Gel purify this fragment!

← 2nd fragment

25

Discussion

30

Just as I suspected DSS4 clone has a mutated EcoRI site as shown by inability of EcoRI to cut out insert (42973-103, lane 2). However, Hind III and Pst I dropped out two fragments. One fragment approximately 400 bp in size will be gel purified and used as a probe (HP400). The 2nd fragment is approximately 80-90 bp and will go into the trash.

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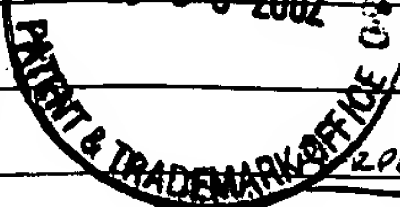
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SUBJECT

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P. 1 purification of clones # 3 and clone # 4

To get purity both clones isolated from Dordic cell substrate
library for use as probes in further exp.

5

Materials & Methods

See protocol from QIAquick Spin Handbook

"QIAquick gel extraction kit" (QIAGEN Cat # 28704)

10

RESULTS

abs 260.0 nm	abs 280.0 nm	bkg abs 320.0 nm	260.0 nm 280.0 nm	280.0 nm 260.0 nm	Protein ug/ml	Nucleic Acid ug/ml
1) 0.0048	0.0027	0.0002	1.8457	0.5418	0.0025	23.1278
2) 0.0109	0.0045	-0.0006	2.2434	0.4457	0.0051	57.5225
3) 0.3912	0.2485	0.0023	1.5793	0.6332	0.2463	1944.6359
4) 0.4535	0.2929	0.0039	1.5558	0.6428	0.2890	2248.1201

$$[C] = \frac{(A_{260} \times (50^{25}/ml) \times D)}{1000} = \frac{0.9}{25}$$

1) DSS4

25

$$(0.0048 \times 50^{25}/ml \times 25) / 1000 = 6^{25}/ml$$

2) DSS02

$$(0.0109 \times 50^{25}/ml \times 25) / 1000 = 13^{25}/ml$$

3) MAIPREP
DSS4

3) DSS4.1

$$(0.3912 \times 50^{25}/ml \times 200) / 1000 = 3.9^{25}/ml$$

4) DSS4.2

$$(0.4535 \times 50^{25}/ml \times 200) / 1000 = 4.5^{25}/ml$$

DISCUSSION:

35

NONE

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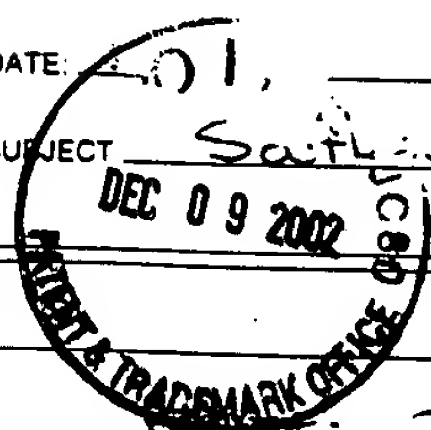
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SUBJECT

South. Hybridization to confirm expression of PSS4 in subtracted library



5

Materials and Methods:

10

10µg of each cDNA was loaded into each well
and run @ 70V for 3 hours.
DNA was stained (see gel photo)

	abs 260.0 nm	abs 280.0 nm	bkg abs 320.0 nm	260.0 nm 280.0 nm	280.0 nm 260.0 nm	Protein ug/ml	Nucleic Acid ug/ml
1)	0.6093	0.4039	0.0198	1.5350	0.6515	0.3840	2947.4448
2)	0.9419	0.5982	0.0306	1.6055	0.6229	0.5676	4556.4517
3)	0.2023	0.1188	0.0005	1.7056	0.5863	0.1183	1008.9186

20

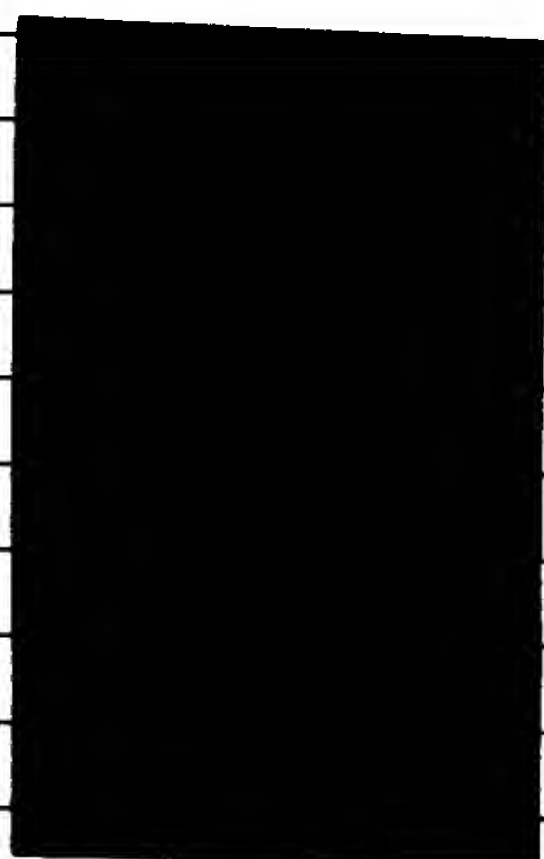
$$[C] = \text{Abs} \times 50^{-1} \text{ml} \times 10 = \%/\text{ml}$$

1) DIZ cDNA

$$0.4039 \times 50^{-1} \text{ml} \times 50 = 1.5 \%/\text{ml}$$

25

2) THPI cDNA $0.9419 \times 50^{-1} \text{ml} \times 50 = 2.4 \%/\text{ml}$

3) DIZ-THPI subtracted cDNA $0.2023 \times 50^{-1} \text{ml} \times 50 =$
c.5%


30

AT Lane 1 : DIZ cDNA
Lane 2 : THPI cDNA
Lane 3 : DIZ-THPI subtracted cDNA

35

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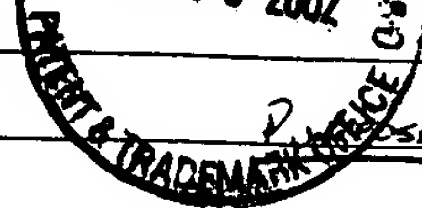
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SUBJECT

3. RACE using DSS4 gene specific primer and QT, QI, QII

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To amplify the remaining 3' end of the unknown DSS4 gene which shows homology to CDB4

5

Materials and Methods

1) cDNA synthesis in Dendritic, THPI, and Jurkat cells

10

poly A⁺ RNA~~DNA~~

2ul

Primer QT(JNF3)

1ul

dH₂O (DEPC)

9ul

12ul

*Incubate @ 70°C for 10 min. Inc 1 min.

15

Add to RNA/primer mix

10x PCR Buffer

2ul

25mM MgCl₂

1ul

10mM dNTP's

2ul

20

0.1M DTT

2ul

7ul

2) Incubate @ 42°C 5 min

Add 1ul Superscript II RT

25

Incubate @ 42°C 1 hour

Stop Rxn by 70°C incubation

Add 1ul RNase H

Store @ -20°C

30

continued 113

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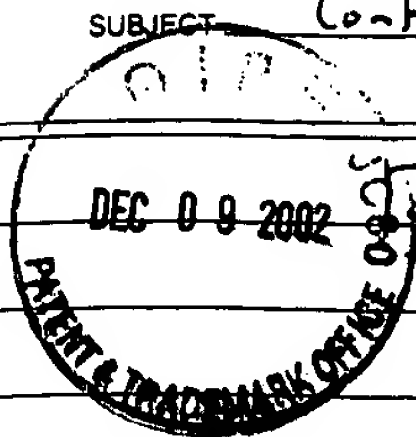
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SUBJECT

Control from 42973-112



Contribution from 42973-112

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2) Primary PCR using JNF1 and Qo(JNF4)

5

JNF1 1 ul

JNF4 1 ul

dNTP's 1 ul

10x PCR Buffer 5 ul

10

~~50x~~ cDNA 1 uldH₂O 39 ul

50x cDNA Amplification mix 1 ul

50 ul

15

PCR parameters: 94°C 1min 1 cycle

94°C 1min

57°C 1min } 25 cycles

72°C 2min

72°C 2min 1 cycle

20

3) Secondary PCR using JNF2 and Qo (JNF5)

JNF2 1 ul

JNF5 1 ul

25

dNTP's 1 ul

10x PCR Buffer 5 ul

1/50 dilution of 1st PCR 1 uldH₂O 39 ul

50x Polymerase mix 1 ul

30

50 ul

PCR parameters: Same as Primary PCR

35

Conforms to 114

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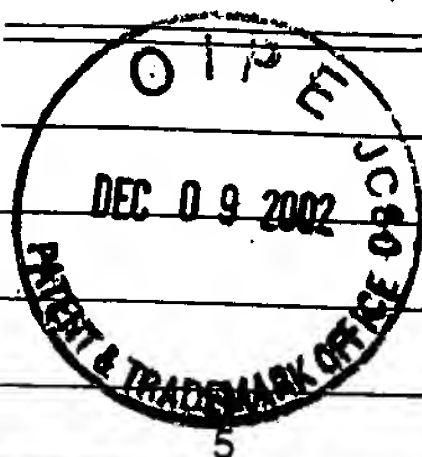
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PROJ. NO.

EXPT. NO.

SUBJECT

Contract for 42423-113



RESULTS

1 2 3 4 5 6 7 8



Lane 1 XH Marker

Lane 2 Dendritic 4-1

Lane 3 THPI 4-1

Lane 4 Jurkat 4-1

Lane 5 Dendritic 2-5

Lane 6 THPI 2-5

Lane 7 Jurkat 2-5

Lane 8 50bp Marker

10

15

Discussion

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25

Lanes 2,3,4 are primary PCR using gsp1(JNF1) and primer Q1(JNF4).
 There seems to be background products. Either extension time of 2 min was
 too long or T_m of annealing was low. Secondary PCR using gsp2(JNF2)
 and primer Q2(JNF5) reduced background considerably, but contains several
 bands which may be nonspecific (Lanes 5,6,7). Continue with RFLP
 experiments, starting with 1/50 dilution of 1st PCR, but increase T_m to
 58°C and reduce extension to 1 min (since most prominent band in
 Jurkat lane is around 900 bp).

END

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EXPT. NO. _____

SUBJECT: _____

Cloning of Full length DCS4 cDNA including 3' UTR.

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10 isolate The full length DCS4 cDNA of size approximately 2.6 kb from 50 μ l of PCR mixture.

5 Materials and MethodsPCRPCR parameters

10	EP (Liu et al, 1997)	2.5 μ l	94°C 1 min
	RP (TJNF4)	1.0 μ l	94°C 1 min
	dNTP's	1.0 μ l	57°C 1 min
	10x PCR Buffer	5.0 μ l	72°C 2.5 min
	Advent Polymerase	1.0 μ l	72°C 1.0 min
	dH ₂ O	38.5 μ l	
	Dendritic cDNA	1.0 μ l	
15		50.0 μ l	

Run all 50 μ l on 1.2% agarose gel.

Isolate 2.5 kb band using gel extraction kit (QIAgen)

Run on 2.0% gel; check conc.

20



Lane 1: 10 marker

Lane 2: 1 μ l isolated PCR fragment

Lane 3: 3 μ l isolated PCR fragment

Lane 4: 2 μ l ϕ marker (1.3 kb band = 16 ng)

Lane 5: 4 μ l ϕ marker (1.3 kb band = 32 ng)

25

Conc. of isolated fragment is too low: ≈ 2 ng/ μ l

Ligate fragment into TA cloning vector

30

Results: Amount of 2.5 kb fragment isolated is very low, but I ligated anyway just to try and isolate a clone.

Discussion: No discussion yet, wait until transformation.

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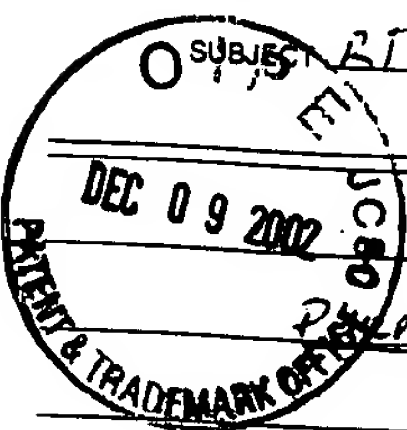
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PROJ. NO. _____

EXPT. NO. _____



SUBJECT: RT-PCR using primers JNF6 and JNF7

Purpose: To optimize parameters for RT-PCR expt., so as to get one band at 600 bp amplified in cell lines which express the DCS4 gene.

5 Materials and Methods

1) 1st PCR: JNF6 primer 1ul cDNA's used:
 JNF7 primer 1ul a) Dendritic
 dNTP's 1ul b) LPS NINT all
 10x PCR Buffer 5ul c) THP1
 Advantage Polymerase 1ul
 dH₂O 40ul
 DNA 1ul
 50ul

Parameters: 94°C 1min
 94°C 1min
 61°C 1min } 30 cycles
 72°C 45sec
 72°C 1min



Lane 1: D10 min
 Lane 2: Dendritic
 Lane 3: LPS NINT
 Lane 4: THP1

RESULT 1: 600 bp band was evident, but a high background was present in reactions. Redo PCR with elevation of annealing temp to 63°C, in order to reduce background.

2) 2nd PCR: JNF6 primer 0.5ul cDNA's used:
 JNF7 primer 0.5ul a) LPS NINT
 dNTP's 0.5ul b) Dendritic
 10x PCR Buffer 2.5ul c) THP1
 Advantage Polymerase 0.5ul
 dH₂O 20.0ul
 cDNA 0.5ul
 25.0ul
 Parameters: 94°C 1min
 94°C 1min
 63°C 1min } 25 cycles
 72°C 45sec
 72°C 1min



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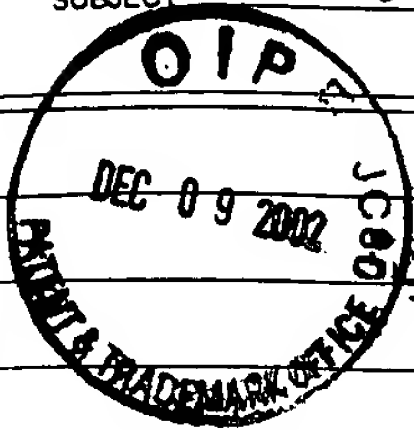
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PROJ. NO. _____

EXPT. NO. _____

SUBJECT

Continued 42973-128



5

Result 2: Significant reduction of background, providing conclusive identification of core binding in LPS Non T cells and no band in THP1 as found earlier on page 42973-125.

10

Discussion: This expt. was done to optimize PCR parameters for RT-PCR of immunological cell lines in order to identify which cell type express the DCS4 transcript. As shown above, LPS Non T cells express DCS4, but THP1 does not. The significance of lack of expression in THP1 mRNA shows the original construction expt up-regulated cDNA's expressed differentially between GM-CSF/IL4 differentiated human monocytes and THP1 monocytes. This is also proven by expression of DCS4 in dendritic cDNA. Repeat PCR for all available cell lines.

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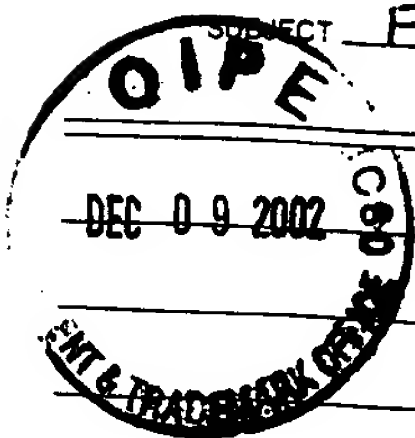
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PROJ. NO.

EXPT. NO.

SUBJECT Full length cDNA sequence and a.a. translation for DCS4



ATGGCTGGTTCCCAACATGCCTCACCTCATCTATATCCTTTGGCAGCTCACAGGGTCA 41
M A G S P T C L T L I Y I L W O L T G S 101
GCAGCCTCTGGACCCGTGAAAGAGCTGGTCCGTTCCGTTGGTGGGGCCGTGACTTTCCCC 161
A A S G P V K E L V G S V G G A V T F P 40
CTGAAGTCCAAAGTAAAGCAAGTTGACTCTATTGTCTGGACCTTCAACACAACCCCTCTT 221
L K S K V K Q V D S I V W T F N T T P L 60
GTCACCATACAGCCAGAAGGGGGCACTATCATAGTGACCCAAATCGTAATAGGGAGAGA 281
V T I Q P E G G T I I V T Q N R N R E R 80
GTAGACTTCCAGATGGAGGCTACTCCCTGAAGCTCAGCAAACTGAAGAAGATGACTCA 341
V D F P D G G Y S L K L S K L K K N D S 100
GGGATCTACTATGTGGGGATATACAGCTCATCACTCCAGCAGCCCTCCACCCAGGAGTAC 401
G I Y Y V G I Y S S S L Q Q P S T Q E Y 120
GTGCTGCATGTCTACGAGCACCTGTCAAAGCCTAAAGTCACCATGGGTCTGCAGAGCAAT 461
V L H V Y E H L S K P K V T M G L Q S N 140
AAGAATGGCACCTGTGTGACCAATCTGACATGCTGCATGGAACATGGGGAAGAGGATGTG 521
K N G T C V T N L T C C M E H G E E D V 160
ATTATACCTGGAAGGCCCTGGGGCAAGCAGCCAAATGAGTCCATAATGGGTCCATCCTC 581
I Y T W K A L G Q A A N E S H N G S I L 180
CCCATCTCCTGGAGATGGGGAGAAAGTGATATGACCTTCATCTGCGTTGCCAGGAACCCT 641
P I S W R W G E S D M T F I C V A R N P 200
GTCAGCAGAACTTCTCAAGCCCCATCCTTGCCAGGAAGCTCTGTGAAGGTGCTGCTGAT 701
V S R N F S S P I L A R K L C E G A A D 220
GACCCAGATTCTCCTCATGGTCCCTCTGTCTCTCTGTTGGTGGCCCTCTGCTCAGTCTC 761
D P D S S M V L L C L L L V P L L L S L 240
TTTGTACTGGGGCTATTTCTTTGGTTTCTGAAGAGAGAGAGACAAGAAGAGTACATTGAA 821
F V L G L F L W F L K R E R Q E E Y I E 260
GAGAAGAAGAGAGTGGACATTTGTCTGGGAAACTCCTAACATATGCCCCATTCTGGAGAG 881
E K K R V D I C R E T P N I C P H S G E 280
AACACAGAGTACGACACAATCCCTCACACTAATAGAACAATCCTAAAGGAAGATCCAGCA 941
N T E Y D T I P H T N R T I L K E D P A 300
AATACGGTTTACTCCACTGTGGAAATACCGAAAAAGATGGAAAATCCCCACTCACTGCTC 1001
N T V Y S T V E I P K K M E N P H S L L 320
ACGATGCCAGACACACCAAGGCTATTTGCCTATGAGAATGTTATCTAGACAGCAGTGCAC 1061
T M P D T P R L F A Y E N V I * 335
TCCCCTAAGTCTCTGCTCAAAAAAAAAACAATTCTCGGCCCAAAGAAAAACAATCAGAAGA 1121
ATTCAGTATTTGACTAGAAACATCAAGGAAGAATGAAGAACGTTGACTTTTTCCAGGA 1181
TAAATTATCTCTGATGCTTCTTTAGATTAAAGAGTTCGTAATTCATCCACTGCTGAGAA 1241
ATCTCCTCAAACCCAGAAGGTTAATCACTTCCATCCCAAAATGGGATTGTGAATGTCAG 1301
CAACCCATAAAAAAGTGCTTAGAAGTATTCCTATAGAAATGTAAATGCAAGGTACACA 1361
TATTAATGACAGCCTGTTGTATTAATGATGGCTCCAGGTGCTGAGTCTGGAGTTTCATTCC 1421
ATCCCAGGGCTTGGATGTGAGGATTATACCAAGAGTCTTGCTACCAGGAGGGCAAGAAGA 1481
CCAAAACAGACAGACAAGTCCAGCAGAAGCAGATGCACCTGACAAAAATGGATGTATTAA 1541
TTGGCTCTATAAACTATGTGCCAGCACTATGCTGAGCTTACACTAATTGGTCAGACGTG 1601
CTGTCTGCCCTCATGAAATTGGCTCCAAATGAATGAATCTTTTATGAGCAGTTGTAGC 1661
AGGCCTGACCACAGATTCCAGAGGGCCAGGTGTGGATCCACAGGACTTGAAGGTCAAAG 1721
TTCACAAAGATGAAGAATCAGGGTAGCTGACCATGTTTGGCAGATACTATAATGGAGACA 1781
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CCCTATTGTAGTAAAAAGTCTTCTTACTATCTTAATAAACAGATATTGTGAGATTCA 2681
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